

Changes in deep-water epibenthic megafaunal assemblages in relation to seabed slope
on the Nigerian margin

Daniel O. B. Jones^{1*}, Charles O. Mrabure² and Andrew R. Gates¹

* corresponding author

¹ SERPENT Project, National Oceanography Centre, European Way, Southampton,
SO14 3ZH, UK.

² Total E & P Nigeria Ltd., 37 Kofo Abayomi Street, Victoria Island, Lagos, Nigeria

Abstract

Local-scale habitat heterogeneity associated with changes in slope is a ubiquitous feature of bathyal continental margins. The response of deep-sea species to high habitat heterogeneity is poorly known and slope can be used as a proxy for many important ecological variables, such as current flow, sedimentation and substratum type. This study determines how slope angle effects megafaunal species density and diversity at the Usan field, offshore Nigeria, between 740-760 m depth. This deep-water area is increasingly exploited for hydrocarbons, yet lacking in baseline biological information. Replicated remotely operated vehicle video transect surveys were carried out using industry infrastructure (through the SERPENT Project) at a representative range of slopes (1°, 3°, 11° and 29°). Twenty-four species of benthic megafaunal invertebrate were found, numerically dominated by the echinoid *Phormosoma placenta*, and nine species of fish were observed. Megafaunal invertebrate deposit feeder density decreased significantly with increasing slope (density range 0.503 - 0.081 individuals m⁻²). Densities of megafaunal suspension feeders were very low except at the highest slope site (mean density 0.17 m⁻²). Overall species richness was greater on steeper slopes, although the richness of deposit feeders was not affected. Reduced labile organic matter in sediments on steeper slopes

likely reduced deposit feeder densities, but increased current flow at higher slopes allowed both increased richness and density of suspension feeders.

Keywords:

Deep-sea; slope; habitat heterogeneity; landscape; *Phormosoma* placenta; Gulf of Guinea, Nigeria; video

1. Introduction

Habitat heterogeneity at local scales is a ubiquitous feature of bathyal continental margins and exerts a fundamental influence on the diversity and structure of benthic communities (Levin and Dayton, 2009). The ‘habitat heterogeneity hypothesis’ is one of the cornerstones of ecology (e.g. MacArthur and Wilson, 1967; Simpson, 1949). It assumes that structurally complex habitats may provide more niches and diverse ways of exploiting the environmental resources and thus increase species diversity (Tews *et al.*, 2004). Characterisation of the seabed in terms of terrain parameters, such as slope and aspect, is a potentially valuable tool for classifying the structural complexity of the seabed and delineating regions of seabed that may support particular fauna and therefore provide a distinct habitat (Wilson *et al.*, 2007). These techniques, broadly classified as habitat mapping, have been successful in shallow waters (Lundblad *et al.*, 2006) and are receiving increased attention in deep water (Jones and Brewer, 2012; Wilson *et al.*, 2007). Slope is a commonly used and important parameter in these analyses, but direct assessment of changes in deep-water assemblages in relation to seabed slope have not (to the author’s knowledge) been assessed quantitatively. Flat areas tend to exhibit different seabed facies and support communities that are different to those inhabiting steeply sloping areas (Lundblad *et al.*, 2006; Schlacher *et al.*, 2007). Slope, as with depth, is not a direct environmental factor that controls biological communities; rather it is a useful proxy for correlated but unmeasured factors. In the case of slope, these environmental factors range across the scale of investigation but include a complex of environmental variables, primarily habitat heterogeneity (Schaff *et al.*, 1992), local current flow (Noble and Mullineaux, 1989; Palardy and Witman, 2011), sedimentation, substratum type, food availability and colonisation (Butman, 1990; Butman *et al.*, 1994; Gage, 1997). Shallow water

research suggests that these factors may lead to increases in density (Sebens, 1985), higher diversity (Palardy and Witman, 2011), elevated growth rates and changes in species distribution (Leichter and Witman, 1997) as well as functional changes in benthic communities, for example favouring suspension feeding over deposit feeding organisms (Jumars and Nowell, 1984). It is important to assess the role of slope-related factors in driving the structure of deep-water benthic faunal assemblages (Levin *et al.*, 2001; Tyler, 1995), particularly in areas of high habitat complexity such as canyons (De Leo *et al.*, 2010; Rowe *et al.*, 1982) and seamounts (Clark *et al.*, 2010; Genin *et al.*, 1986).

By combining spatially-accurate quantitative seabed imaging techniques with high-resolution acoustic data on submarine topography and sediment properties fine-scale patterns in megafaunal biology can be linked to the broad-scale patterns in habitat type and subsea landscape (Baguley *et al.*, 2006; Wilson *et al.*, 2007). Given the sheer size and difficulties of accessing the deep-sea environment, accurate extrapolation of fine-scale observations using information on the important physical controls for faunal distribution is vital to describe the important broad-scale patterns in benthic biology (Sanchez *et al.*, 2008; Williams *et al.*, 2010). Characterising variation in biodiversity and biological assemblage structure in response to specific terrain parameters, such as slope, are important to help understand the underlying mechanisms behind these patterns.

The megabenthic communities of the offshore waters of the Gulf of Guinea region are poorly known (Le Loeuff, 1993; Le Loeuff and von Cosel, 1998; Lebrato and Jones, 2009) compared to the Canary region to the north (Duineveld *et al.*, 1993a; Duineveld *et al.*, 1993b; Henriques *et al.*, 2002; Jones and Brewer, 2012; Keller and Pasternak, 2002; Merrett and Domanski, 1985; Merrett and Marshall, 1981). The Benguela region to the south of the Gulf of Guinea has also received some attention, particularly on the shelf and slope (Roeleveld *et al.*, 1992; Roy *et al.*, 2007; Uriz, 1988) as well as the abyss (Kroncke and Turkay, 2003; Levin and Gooday, 2001; Saiz-Salinas, 2007; Thandar, 1999; Vinogradova *et al.*, 1990; Zibrowius and Gili, 1990). The Gulf of Guinea region is heavily exploited for natural resources, primarily hydrocarbon resources (Zabanbark, 2002) and fish (FAO Fishery Committee for the Eastern Central Atlantic, 1991) and is subject to high levels of anthropogenic impact

from pollution, mostly of terrestrial origin (Scheren *et al.*, 2002). The limited environmental information is mostly collected by oil and gas companies and not available in the scientific literature. However, it is crucial that baseline quantitative environmental information is available for successful management of this increasingly exploited ecosystem.

This paper provides the first description of deep-water megabenthic assemblages at a deep-water Nigerian site. Quantitative remotely operated vehicle (ROV) imaging allowed the effect of various seabed slopes on assemblage density, composition and diversity to be assessed. Specifically, the following hypotheses were tested: 1) increases in environmental heterogeneity associated with slope will increase megafaunal species richness, 2) increased slope will favour suspension feeding lifestyles at the expense of deposit feeding and hence change the relative abundances of these functional groups and 3) the combination of these effects will lead to differences in megafaunal assemblage structure between habitats with biologically significant variation in environmental conditions resulting from changes in slope.

2. Methods

2.1. Study Site

This study focuses on a knoll at the west of the Usan field (within Nigerian offshore oil prospecting license block 222), an area that will be developed for hydrocarbon production, located on the Nigerian continental margin in the Gulf of Guinea.

Environmental baseline surveys (Total E. & P. Nigeria Ltd., 2007) in the main Usan field (centred 03° 31.800' N 007° 26.400' E; slope < 1°) collected boxcore samples of seabed sediments. The silty clay sediment was comprised of silt (50.1 ± 1.1 %; mean \pm standard deviation), clay (34.9 ± 2.0 %) and sand (15.1 ± 1.0 %), with an organic content of 5.8 ± 1.3 % and a total organic carbon content of 3.3 ± 0.7 %. Sediments contained abundant material of terrigenous origin (e.g. seeds, pollen, plant material). Sediments supported a macrofaunal (>0.5 mm) assemblage with low density (69 ± 53 individuals m⁻²) and diversity (57 species). The macrofauna was dominated by molluscs (38.6%), arthropods (22.0%), protozoans (17.6%) and annelids (16.3%),

with the remainder comprised of echinoderms, cnidarians, poriferans and hemichordates (Total E. & P. Nigeria Ltd., 2007).

Shipboard measurements of seawater collected near the seabed at Usan had dissolved oxygen levels (mean = 5.84 mg L⁻¹ [54.7 % saturation], range = 4.41 [40.8 %] to 9.54 [88.4 %] mg L⁻¹) that were not hypoxic (dissolved oxygen is < 0.7 mg L⁻¹ (Helly and Levin, 2004)). *In situ* physical measurements of the near-bottom seawater (at 750 ± 10 m water depth) made during this study showed low temperatures (5.32 ± 0.07 °C; mean ± standard deviation) and consistent salinities (34.76 ± 0.01).

2.2. Survey Design

Four habitats, each with different slope conditions (1°, 3°, 11° and 29°), were assessed at Usan (Figure 1) through the SERPENT project (Jones, 2009). The slope conditions were labelled 1°, 3°, 11° and 29°. For each slope condition an area was chosen within the Usan field using detailed bathymetric charts (Figure 1) that fulfilled the following criteria: constant slope, a depth between 740 and 760 m and a distance of at least 1 km from any drilling activity to ensure no effect of disturbance (Jones *et al.*, 2006). Three replicate remotely operated vehicle (ROV) video transects (200 m length) were performed at random locations within each slope condition (S1). Survey start points and survey direction were identified using a random number generator in Microsoft Excel. Transects were examined by plotting them onto the chart. Any transect that extended beyond the boundaries of identified areas was discarded and another generated using the same method. In the area of highest slope headings were constrained between 350 to 360°, 0 to 10° and 170 to 190° to maintain the depth range of transects but otherwise were random. Each transect was regarded as a sampling unit (i.e. all organism densities were totalled for each 200 m transect) and used in all subsequent analysis. Three replicate transects at four slope levels resulted in a total of 12 sampling stations. All positional information was recorded in Universal Transverse Mercator (UTM) zone 32 N based on the World Geodetic Society 1984 datum.

2.3. Data collection

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170 Data were collected using an Oceaneering Millennium ROV equipped with a colour
171 video camera (Insite Pacific Pegasus), sonar and ultra-short baseline navigation
172 transponder. Cameras were mounted on a pan and tilt unit at the front of the ROV,
173 which enabled oblique video to be taken. Before each transect the video was zoomed
174 out to maximum extent and the camera was set to its most vertical angle (30° below
175 the horizontal).

176

177 In every transect the ROV was run in a straight line, on a set bearing and at a constant
178 speed ($\sim 0.3 \text{ m s}^{-1}$). Vehicle altitude (distance from seabed to the base of the ROV)
179 was kept constant at 1 m, which equated to a 143 cm camera altitude (from the centre
180 of the camera). Transect width (mean of 1.72 m; max variation $\pm 0.2 \text{ m}$) was
181 calculated from the camera acceptance angles (Jones *et al.*, 2006). The camera
182 horizontal acceptance angle (at full wide angle) was 55° and the vertical acceptance
183 angle 43°. Transect width was also verified by passing over objects of known size on
184 the seafloor. The optical resolution of the cameras permitted all organisms larger than
185 50 mm to be reliably distinguished. In addition to the transects, previous ROV suction
186 sampling at the nearby Akpo site (03°06.950' N 006°49.120' E) and a separate survey
187 to carry out detailed video inspection of selected individual organisms aided species
188 identifications (Jones *et al.*, 2006).

189

190 2.4. Data Analysis

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192 Video transects were replayed at half speed and all visible organisms were counted
193 along the entire transect. Fauna were identified to the highest taxonomic resolution
194 possible, in some cases to species. Megafauna that could not be identified to species
195 were grouped into morphologically distinguishable entities (referred to as taxa). It
196 should be emphasised that this approach may underestimate the number of species
197 present. Colonial organisms were counted as single individuals. Infaunal species were
198 counted if enough of their body was visible for identification. Benthic fish were
199 recorded but not included in subsequent quantitative analysis. Megafauna were
200 classified into deposit and suspension feeders based on the literature descriptions.

201

202 Four indices of diversity were calculated in order to assess both the dominance and
203 species richness aspects of diversity (Magurran 2003, Gotelli & Colwell 2001). These
204 were the number of taxa in a transect or group of transects (a simple measure of
205 species richness, S), rarefied species richness (estimated species in 60 individuals:
206 $ES_{(60)}$), evenness (as measured by Pielou's evenness, J') and Shannon-Wiener species
207 diversity (H' base log e). Numbers of organisms were expressed as abundances
208 (numbers in each sampling unit) or as densities (no. m^{-2}). Diversity indices were
209 calculated using PRIMER v6 (Clarke and Warwick, 2001).

210

211 The structure of the megabenthic assemblage was visualised using multidimensional
212 scaling plots and sites compared using multivariate community analysis, specifically
213 PERMANOVA (Anderson, 2001) and Multivariate dispersion analysis (Anderson,
214 2006). Multivariate community analysis was based on abundances of all taxa, a fourth
215 root transformation being applied to buffer the influence of dominant taxa (Field *et*
216 *al.*, 1982). Similarities were calculated using Bray-Curtis coefficients (Bray and
217 Curtis, 1957). The similarity values were subjected to both classification (hierarchical
218 group-average clustering) and ordination (non-metric multi-dimensional scaling)
219 using the software PRIMER v6 (Clarke and Warwick, 2001). PERMANOVA and
220 Multivariate dispersion analysis were carried out using the 'adonis' and 'betadisper'
221 functions respectively using the R package Vegan (Oksanen *et al.*, 2011) in the R
222 programming environment (R Development Core Team, 2010).

223

224 Data were analysed following one-way analysis of variance (ANOVA) design with
225 slope as a fixed factor. This was implemented in the univariate case by generalised
226 linear models and in the multivariate case by one-way PERMANOVA (Anderson,
227 2001). Independent generalised linear models were developed (Dobson and Barnett,
228 2008) to examine whether the taxon richness (number of species in a transect) and
229 density (as numbers of epibenthic invertebrates in a transect) of both suspension and
230 deposit feeding taxa could be explained by the seabed slope (treated as categorical
231 data). The residual deviance for the count data (48.44 for suspension feeders; 97.26
232 for deposit feeders) was greater than the residual degrees of freedom (8 for both),

indicating overdispersion. The models were, therefore, fitted with quasi-Poisson errors. The models for species richness, evenness and species diversity were fitted with Gaussian errors. Models were fitted using the R function glm and the ANOVA function of the R package CAR (companion to applied regression) (Fox and Weisberg, 2011).

3. Results

The seabed was almost entirely composed of fine grain-sized sediment, no individual grains were visible in video (Figure 2), even when fully zoomed in. This suggested that the largest possible grain size is fine sand (< 0.5 mm) but observations of resuspension indicate that it was finer and classified as mud (< 125 μ m). Hard substrata, composed of igneous rock (identified from samples), were visible within the areas of highest slope. These hard surfaces provided habitat for numerous specialist taxa (Table 1).

3.1. Nature of the benthic community

A total of 24 invertebrate megafaunal taxa representing six phyla were observed in the video transects at Usan (Table 1, Figure 3). In terms of species richness and numerical abundance, echinoderms were most abundant: 64% of the fauna were accounted for by 10 echinoderm taxa. Although eight cnidarian taxa were identified, they only represented 4% of the total density. Crustaceans were numerically important (31% total fauna) but only represented by three taxa, one of which, a galatheid, accounted for the vast majority of individuals. Of lesser importance were the poriferans, molluscs and enteropneusts, all with only one observed taxon and together representing $<1\%$ of the total invertebrates.

The regular echinothurid echinoid *Phormosoma placenta* was overwhelmingly dominant, particularly in the flatter areas, accounting for 53% of the invertebrates observed. Two other taxa were numerically important; the galatheid *Munidopsis* sp.

represented 29% of megafauna and an unidentified large ophiuroid represented 8%.
The other taxa observed were each of low abundance (<3% of megafauna).

Although not considered further in this analysis, fish were observed at all the sites (S2). A small, dark-coloured, midwater gonostomatid fish was most abundant representing over 47 % of the fish fauna. The true benthic lophiiform *Dibranchius* sp. was common and represented 27% of the fish. The other major taxon was the nettastomatid anguilliform *Nettastoma melanura* which accounted for 18% of the fish. The demersal grenadier *Corphaenoides* sp. and the morid gadiform *Laenomena laureysi*? each represented around 2% of the fish seen in the video records.

3.2. Variation between habitats

3.2.1. Faunal density

Faunal density varied between habitats of different slope, with lower total densities in areas of higher slope. When total density is split into more ecologically meaningful units, based on feeding mode, the trends were even clearer (Figure 4). Deposit feeders made up the majority of the megafauna observed; however, when analysed alone, densities of deposit feeders decreased significantly with greater slopes (L-ratio = 16.09, d.f. = 3, $p < 0.01$). Densities of suspension feeders were very low except at the habitat with the highest slope; even here the density was less than half (0.17 m^{-2}) that of deposit feeders (0.40 m^{-2}). The increase in deposit feeder densities with slope was significant (L-ratio = 35.41, d.f. = 3, $p < 0.001$).

Of the major taxa, *Phormosoma placenta* had higher densities at low slope sites (L-ratio = 16.11, d.f. = 3, $p < 0.01$) as did the galatheid *Munidopsis* sp. (L-ratio = 88.37, d.f. = 3, $p < 0.001$). Ophiuroid densities were significantly higher in areas of higher slope (L-ratio = 24.88, d.f. = 3, $p < 0.001$).

3.2.2. Megafaunal diversity and assemblage structure

The slope-related differences in the habitat had a significant effect on the number of taxa found (Figure 4; L-ratio = 9.33, d.f. = 3, $p < 0.05$). Pairwise tests, however, only revealed significant differences between the habitat with the lowest slope (1°) and the highest slope (29°). The principal difference in the diversity of sites was caused by the presence of obligate epilithic species at the sites where hard substratum was exposed (see Table 1). In soft sediment areas there was some variation in species complement resulting from the apparently random presence or absence of rare species.

Species diversity indices (Figure 4) revealed that the species richness element of diversity ($ES_{(60)}$: L-ratio = 35.42, d.f. = 3, $p < 0.001$) and the evenness component (J' L-ratio = 66.14, d.f. = 3, $p < 0.001$) showed a highly significant positive linear relationship with increasing slope of the habitat. As would be expected, a combined index showed similar patterns with slope (H' ; Figure 4). There were significant pairwise differences between all the habitats ($p < 0.05$) in $ES_{(60)}$, J' and H' except between 1 and 3 degrees in the combined Shannon-Wiener Index ($p = ns$).

Overall, multivariate community similarity was significantly different between habitats of different slope (PERMANOVA: $F = 9.55$, d.f. = 1, 11, $p < 0.001$; Figure 5). Multivariate community similarity was high ($> 60\%$) and between-site dispersion low between the habitats with lower slope (1 , 3 and 11°), whereas the habitat with highest slope (29°) was more heterogeneous (Multivariate dispersion analysis: average distance to centroid 0.08 for 1° ; 0.14 for 3° ; 0.16 for 11° and 0.30 for 29° ; differences not significant in permutation test $F = 2.57$, d.f. = 3, $p = 0.13$). Two sites (29B and 29C) formed a distinct group (with 77.5 % similarity) and one site (29A) was intermediate between the highest and lower slope sites, having slightly higher affinity with the lower slope sites (similarity 58.8%) than with the higher slope sites (similarity 35.9%).

4. Discussion

4.1. General characteristics of the megafaunal assemblages at Usan

Knowledge of West African megafauna is limited, making it difficult to compare the assemblage of invertebrates found at Usan to those at other locations. However, the common species are known and the limited evidence suggests they are abundant off West Africa. *Phormosoma placenta* is a common, often dominant, species on both sides of the Atlantic (Jones and Brewer, 2012; Laguarda-Figueras *et al.*, 2005; Sanchez *et al.*, 2008). The large ophiroid observed at Usan is very similar to a species found off Mauritania (Jones and Brewer, 2012), Ivory Coast and Angola (author's observation; <http://archive.serpentproject.com/151/>), while a galatheid resembling the species observed in the present study has been seen in images obtained off Ivory Coast (Lebrato and Jones, 2009). It should be noted that even generic differences are not always evident in photographs. The rarer species quantified in our survey, and those seen outside the transect areas, likely include new species. However, standard-resolution video and image data in general is not usually sufficient to resolve the details necessary for species-level taxonomy.

The density of the megafaunal assemblages at Usan seems to be low, but in the range of those found elsewhere (Figure 6). The nearest comparative area of an equivalent depth is located off Ivory Coast where megafaunal densities were over three times greater than at our study site. However, the Ivory Coast site experienced significantly enhanced nutrient inputs from pyrosome carcasses (Lebrato and Jones, 2009). Few other data are available for the East Atlantic at equivalent depths. Studies conducted in deeper areas reported a range of megafaunal densities (Figure 6) that are usually lower than at the Usan site. Of the deeper studies, only those areas that underlie highly productive surface waters, for example the eutrophic EUMELI site off Mauritania (Galeron *et al.*, 2000), have yielded comparable values.

4.2. Environmental consequences of differences in slope

Although four species were present at habitats with all the levels of slope investigated, ten of the taxa found at the habitat with the highest slope (29) were confined to this habitat. This habitat had exposed hard substrata and hence a unique community of epilithic megafauna. Most of the epilithic organisms were not found on soft sediments and many, for example gorgonians, had obvious morphological adaptations for attachment to rocky surfaces. Hard substratum communities are probably rare on the West African margin, having only been described off Mauritania where they are associated with corals (Colman *et al.*, 2005). Other studies of hard substrata near west Africa are on the mid-Atlantic ridge at non-vent (Gebruk *et al.*, 2010) and vent sites (Van Dover *et al.*, 2002), volcanic island groups, for example the Azores, or seamounts (Hall-Spencer *et al.*, 2007). The megafaunal assemblage found on hard substrata at Usan superficially resembles that found at non-vent hard substrata in the northern tropical Atlantic (in terms of faunal groups present) and from hard substrata in the deep sea in general (e.g. Lundsten *et al.*, 2009). However, without accurate species-level identifications either here or in the megafaunal literature generally, further patterns are difficult to assess.

The large reductions in density of deposit-feeding species at Usan with increases in habitat slope likely reflect reductions in food availability for deposit feeders. These patterns are unlikely to be related to the presence of hard substratum at the steepest slope site. Slope is expected to affect food availability in two ways. Firstly, increased slope will probably increase the lateral advection of food particles, reducing their residence times and hence availability for deposit feeders. Decreases in food availability have been repeatedly shown to be responsible for reductions in benthic megafaunal density in the deep sea (Galeron *et al.*, 2000; Thurston *et al.*, 1994). Secondly, increases in slope are likely to enhance local current speeds and hence lead to reductions in the deposition and increases in the winnowing of fine, nutritious particles (Levin *et al.*, 2001). Increased flow favours suspension feeding organisms over deposit feeders (Jumars and Nowell, 1984) and actively enhances recruitment, driving increased species density and elevated species richness (Palardy and Witman, 2011).

Increases in slope were associated with increased overall megafaunal diversity at Usan despite a decrease in the number of deposit-feeding species in the habitat with the steepest slopes. There was slightly (although not significantly) higher diversity, and consistent changes in multivariate community composition, in areas of steep slope without hard substrata. In the absence of other factors, reduced food availability tends to lead to reductions in diversity (Levin *et al.*, 2001; Waide *et al.*, 1999), as demonstrated in the tropical Atlantic deep sea (Cosson *et al.*, 1997). Organic particles are more likely to settle and become available for deposit feeders at low slopes (Fontanier *et al.*, 2008). At high slopes resuspension and current-mediated lateral transport of organic material is likely (Schlacher *et al.*, 2007; Van Gaever *et al.*, 2009), which may favour a suspension-feeding lifestyle. The trends in diversity observed at Usan are consistent with predictions based on food availability. Higher levels of organic matter may result in reduced diversity in areas, such as oxygen minimum zones, where the organic inputs are already high and associated with oxygen depletion (Levin *et al.*, 2001), which was not observed at Usan. In addition to influencing the organic content of the sediment, increases in slope are also likely to affect the substratum heterogeneity. Sediment grain size will likely become coarser and potentially more heterogeneous, increasing niche availability and hence deposit feeder diversity (Etter and Grassle, 1992; Wheatcroft, 1992), although this was not clear in the present study. In the Usan area the sites with steep slope have exposed hard substrata, presumably reflecting limited deposition. This increase in habitat heterogeneity is responsible for the clear and significant increases in diversity and changes in multivariate community composition at the steepest slope sites. An entirely different range of animals is able to colonise hard substrata and hence, where this is present, diversity is likely to increase (Levin *et al.*, 2001).

Deep-sea biology requires more detailed local-scale studies of faunal distribution in relation to habitat features. Physical gradients in biological assemblages associated with slope occur in many ecosystems (de Castilho *et al.*, 2006); this study extends these findings to the African deep sea. Considerable areas of the ocean margins have features of high topography and sloping seabed, such as described here. This study has shown clear reductions in deposit-feeding megafaunal density with slope. If this trend is typical for deep-water habitats, it should be incorporated into model estimates

of megafaunal densities, particularly on areas of elevated topography. Diversity of benthic megafauna does not display such a clear relationship with slope and is likely to be controlled by site-specific responses to changes in habitat heterogeneity and food availability.

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Table 1: Invertebrate megafauna densities (no m⁻²) at each transect at Usan. Slope is represented by numbers: 1 = 1°, 3 = 3°, 11 = 11°, 29 = 29°. Site letters represent replicates. Asterisk taxa (*) are known to prefer hard substrata and double asterisk (**) only live on hard substrata (obligately epilithic). Deposit feeders (D) and suspension feeders (S) are labelled. See figure 3 for images of the taxa.

	1A	1B	1C	3A	3B	3C	11A	11B	11C	29A	29B	29C
Hexactinellid**S	0	0	0	0	0	0	0	0	0	0	0.006	0.003
White antipatharian**S	0	0	0	0	0	0	0	0	0	0	0.047	0.061
Whip-like cnidarian (sand) S	0	0	0	0	0.003	0	0.003	0.003	0.003	0	0	0
Whip-like cnidarian (rock, white)** S	0	0	0	0	0	0	0	0	0	0	0.009	0.009
<i>Pennatulaea</i> sp. S	0	0	0	0	0	0	0	0	0	0	0.006	0
Gorgonian fan** S	0	0	0	0	0	0	0	0	0	0	0	0.009
<i>Actinoscypha</i> sp.	0.003	0	0.003	0	0	0	0	0	0	0	0	0
Edwardsiid anemone	0	0	0.003	0	0	0	0	0	0	0	0	0
Striped anemone	0	0	0.003	0.003	0	0.003	0	0	0	0	0	0
<i>Phormosoma placenta</i> D	0.343	0.395	0.302	0.140	0.233	0.352	0.201	0.009	0.012	0.090	0.023	0.081
Red echinoid	0	0	0	0	0	0	0	0	0	0.003	0.012	0.003
White echinoid	0	0	0	0	0	0	0	0	0	0.003	0	0.003
White <i>Diadema</i> sp.	0	0	0	0	0	0	0	0	0	0.003	0.009	0.006
Ophiuroid D	0.012	0	0	0.009	0.020	0.032	0.064	0.070	0.038	0.029	0.052	0.026
Brisingid* S	0	0	0	0	0	0	0	0	0	0	0.012	0.003
Holothurian D	0.012	0.003	0	0	0.003	0.003	0	0	0	0	0	0
<i>Mesothuria</i> sp. D	0	0.009	0.006	0.003	0	0	0.003	0.003	0.017	0	0	0
<i>Benthothuria</i> sp. D	0	0	0	0.006	0	0	0	0	0.003	0.003	0	0
Commatalid* S	0	0	0	0	0	0	0	0	0	0	0.003	0
Galatheid	0.090	0.070	0.047	0.142	0.180	0.116	0.180	0.172	0.172	0.032	0.006	0.015
Prawn	0.003	0.009	0.015	0.003	0.009	0.003	0.006	0.003	0.009	0.012	0	0
c.f. <i>Nephrops</i> sp.?	0	0	0	0.003	0.003	0	0.003	0.003	0	0	0	0
Opisthobranch D	0	0	0	0.003	0	0	0	0	0	0	0	0
Enteropneust D	0	0	0	0	0	0	0	0	0.003	0	0	0
TOTAL	0.462	0.485	0.378	0.311	0.451	0.509	0.459	0.262	0.256	0.174	0.183	0.218
Total suspension feeders	0	0	0	0	0.003	0	0.003	0.003	0.003	0	0.081	0.084
Total deposit feeders	0.456	0.477	0.355	0.302	0.436	0.503	0.448	0.253	0.244	0.154	0.081	0.122

FIGURES

Figure 1: Bathymetry of Usan area showing the position of ROV seabed transects. The lower panel shows the higher slope transects in more detail. The inset on the upper panel shows the position of the study site in the Gulf of Guinea. The length of the 200 m long ROV transects is drawn to scale.

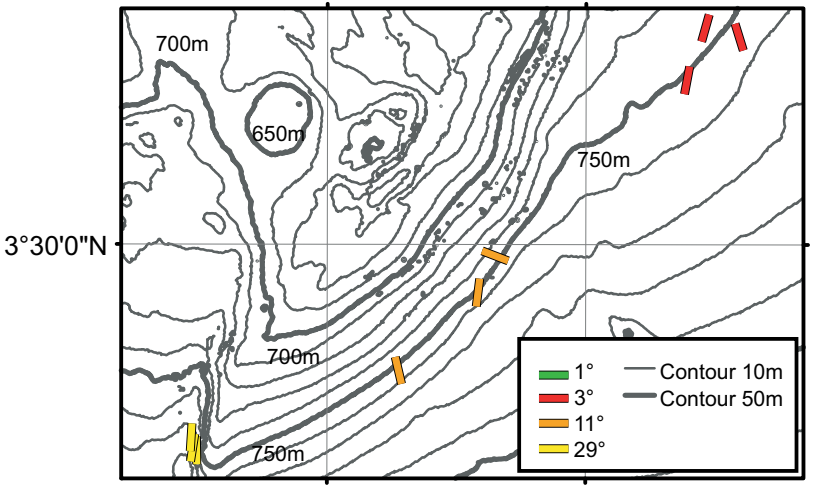
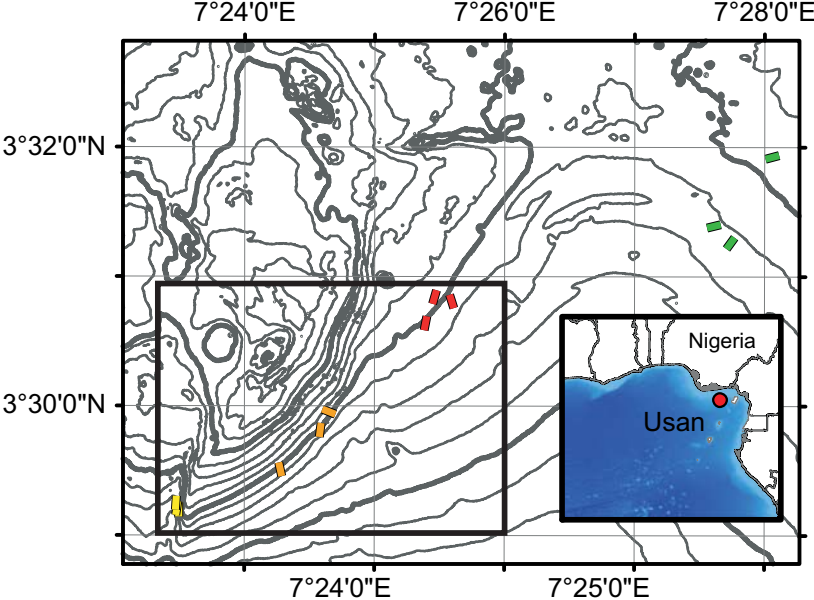
Figure 2: Example pictures of seabed habitats with different slope found at Usan: A) 1° slope, B) 3° slope, C) 11° slope, D) soft-substratum area with 29° slope and E) hard-substratum area with 29° slope. Images A to D are images from the video transects cropped to show an approximately 0.5 m wide section of seafloor at the base of the image. Image E is a more oblique image cropped to show an approximately 1 m wide section of seafloor at the base of the image.

Figure 3: Examples of megafauna found at Usan: A) Hexactinellid (~150 mm diameter), B) White antipatharian (~200 mm height), C) Whip-like cnidarian (found on rocks, ~ 400 mm height), D) Whip-like cnidarian (found on sand, ~ 300 mm height), E) Gorgonian fan (~ 1m height), F) *Pennatulaea* sp. (~300 mm height), G) Brisingid (arm length ~ 250mm), H) *Phormosoma placenta* (50-120 mm diameter), I) Red echinoid (70 mm diameter), J) White *Diadema* sp. (~150 mm diameter including spines), K) Ophiuroid (~150 mm arm-tip to arm-tip), L) Galatheid (up to 100 mm length), M) *Prawn (~100 mm total length), N) *Cerianthid? (~ 100 mm total diameter), O & P) *Lithodid crab (c.f. *Paralomis cristulata*, ~100 carapace width). * species observed but not seen in quantitative transects.

Figure 4: Response of megafaunal assemblage indices to slope at the Usan area, offshore Nigeria. A) Megafaunal invertebrate deposit (hollow circles) and suspension feeding (filled circles) assemblage total densities. B) Number of megafaunal invertebrate deposit (hollow circles) and suspension feeding (filled circles) taxa. C) Evenness of entire invertebrate megafaunal assemblage measured by Pielou's evenness (J'). D) Shannon-Wiener (log base e) combined diversity index (H') of entire invertebrate megafaunal assemblage. Errors bars on all graphs represent standard deviations and points represent the mean density of 3 replicates. For A) and B) the megafauna was separated into deposit (hollow circles) and suspension (filled circles) feeders.

Figure 5: Multidimensional Scaling Ordination of Bray-Curtis similarities (4th root transformed) of Usan Invertebrate megafaunal assemblage. Overlaid are similarity levels from hierarchical cluster analysis (as percentages). Slope is represented by numbers equating to the slope in degrees. Letters represent replicates.

Figure 6: Megafaunal density (no hectare⁻¹) compared with depth for the Eastern Atlantic. East Atlantic literature data from Bay of Biscay south to Angola (Feldt *et al.*, 1989; Galeron *et al.*, 2000; Kroncke and Turkay, 2003; Lebrato and Jones, 2009; Sibuet *et al.*, 1989; Sibuet *et al.*, 1984; Sibuet and Segonzac, 1985; Thurston *et al.*, 1994; Tietjen, 1992)



A



B



C

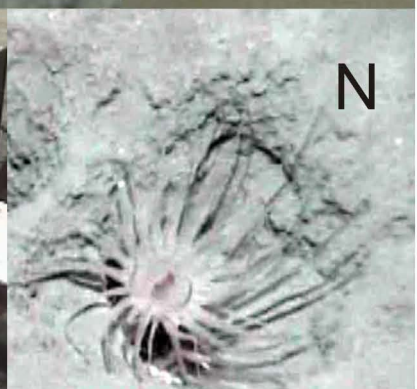
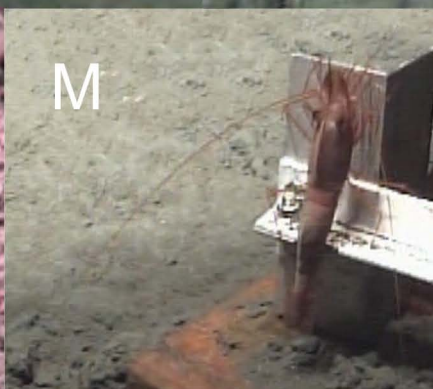
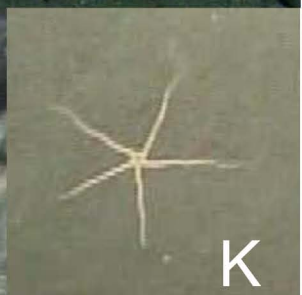
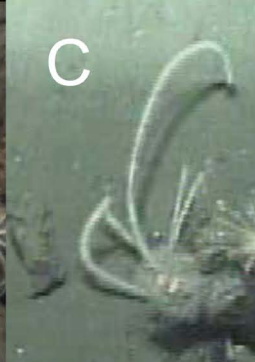


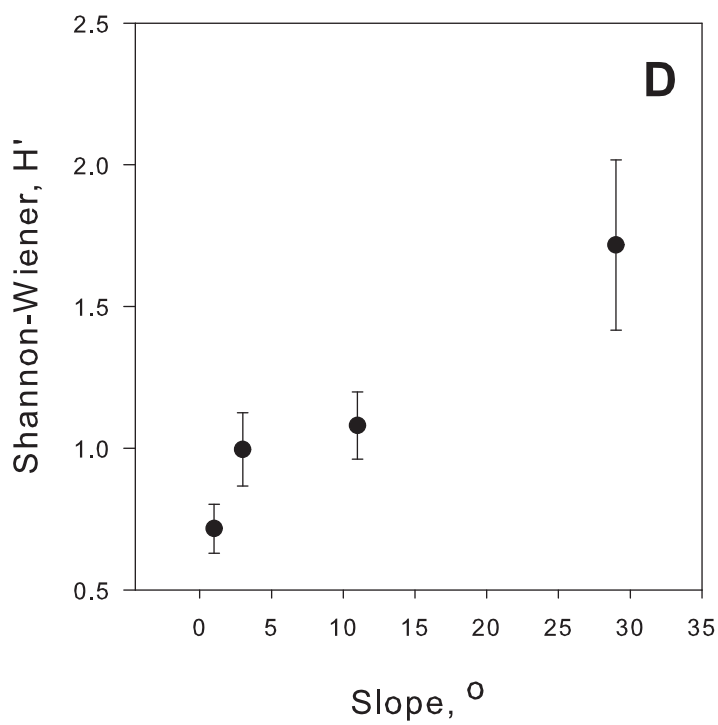
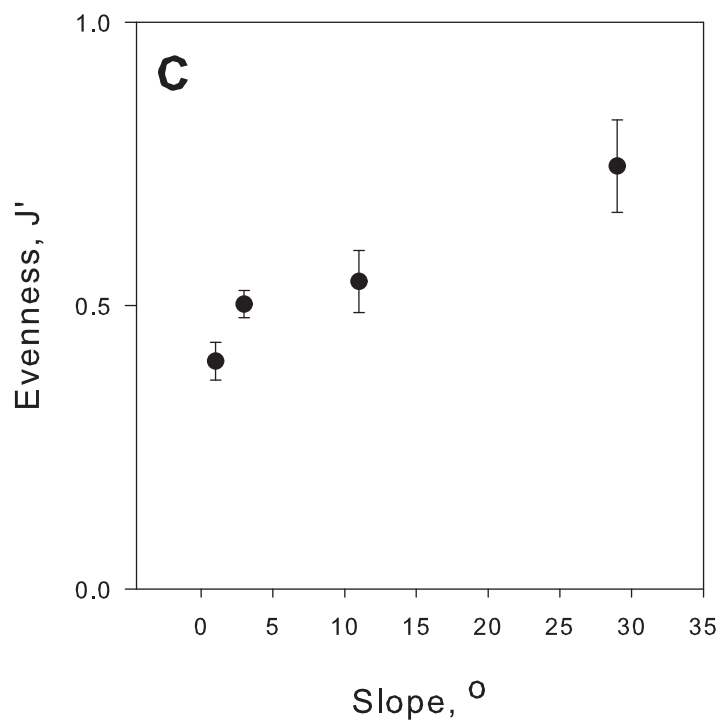
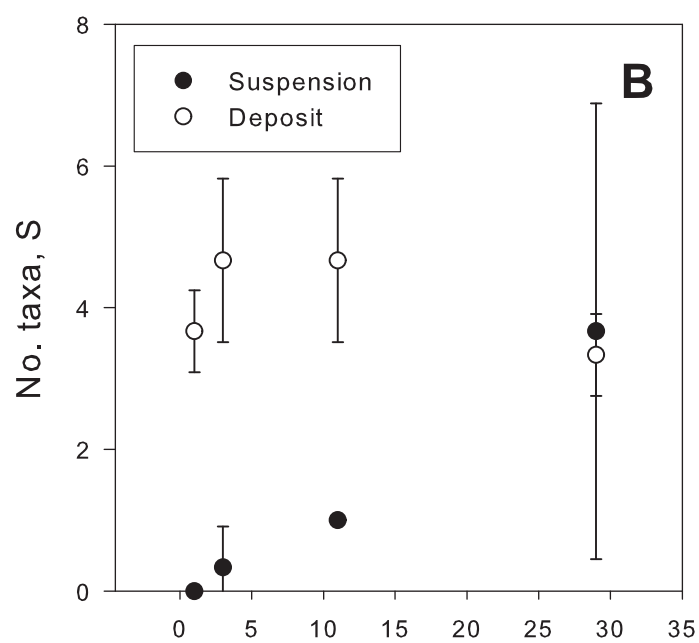
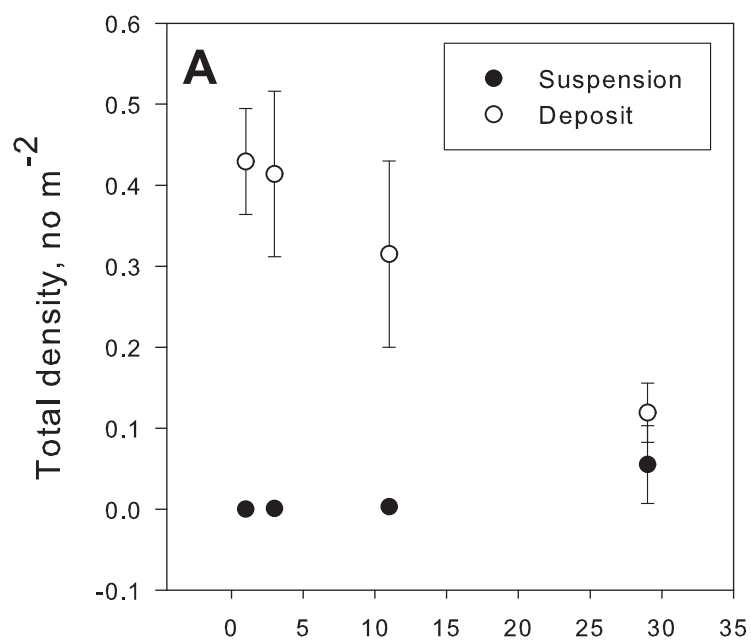
D



E







2D Stress: 0.06

